	•
je sjês	
1000	
586- \$189	
##	
7 3 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
ija vila	

g protein-2 down (0 up - 16 down) down (4 up -12 down) down (1-15) nc (13-13) receptor alpha down (0-13) nc (8-10) down (0-13) down (1-13) down (1-13) nc (9-10) down (1-12) nc (3-3) down (1-12) nc (3-3)	Datenbank-Nr. Names Personal Inc.	Vergleich Endometriose wir in der Schauff in der Sc	Verglejch/Endometriose Versus/Normali(prolitiphase)	Vergleich sekreversus prolipphase
binding protein-2 down (1-15) nc (13-13) u receptor precursor down (0-14) nc (6-2) actor receptor alpha down (0-13) down (0-8) 1 (COL.18A1) down (1-13) down (4-13) 3 down (1-13) nc (9-10) 4 down (0-12) nc (0-0) 1 chain down (1-12) nc (3-3)		down (0 up - 16 down)	dn	up (18 up - 1 down)
rsor down (0-14) nc (6-2) pha down (0-13) nc (8-10) down (1-13) down (4-13) down (1-13) nc (9-10) down (0-12) nc (9-10) down (0-12) nc (9-10) down (1-12) nc (9-10)	X02761, fibronectin (FN precuisor)	down (1-15)	nc (13-13)	up (17-2)
RA) down (0-13) nc (8-10) L RA) down factor receptor alpha down (0-13) down (0-8) fown (0-13) 2, subtilisin-like protein (PACE4) down (1-13) down (4-13) nc (9-10) 3, laminin M chain (merosin) down (0-12) nc (0-0) 6, U77846, Elastin down (0-12) nc (3-3) 0, type IV collagen alpha -2 chain down (1-12) nc (9-10)	် က	down (0-14)	nc (6-2)	(1-6) dn
down (0-13) nc (8-10) down (0-13) down (0-8) down (1-13) nc (9-10) down (0-12) nc (0-0) down (0-12) nc (3-3) down (1-12) nc (9-10)	(PTK7)			(17.0)
outsilisin-like protein (PACE4) down (0-13) down (4-13) to (9-10) ubtilisin-like protein (PACE4) down (1-13) nc (9-10) iminin M chain (merosin) down (1-13) nc (0-0) J77846, Elastin down (0-12) nc (0-0) ype IV collagen alpha -2 chain down (1-12) nc (3-3) down (1-12) nc (9-10)	M21574, platelet-derived growth factor receptor alpha	down (0-13)	nc (8-10)	(o m) dn
collagen type XVIII alpha 1 (COL.18A1) down (0-13) down (0-8) subtilisin-like protein (PACE4) down (1-13) nc (9-10) laminin M chain (merosin) down (1-13) nc (9-10) , U77846, Elastin down (0-12) nc (3-3) , type IV collagen alpha -2 chain down (1-12) nc (9-10)	(PDGFRA)			(47.0)
subtilisin-like protein (PACE4) down (1-13) down (4-13) laminin M chain (merosin) down (1-13) nc (9-10) , U77846, Elastin down (0-12) nc (0-0) , type IV collagen alpha -2 chain down (1-12) nc (3-3) down (1-12) nc (9-10)	1 22548, collagen type XVIII alpha 1 (COL18A1)	down (0-13)	down (0-8)	(o-11) dn
down (1-13) nc (9-10) down (0-12) nc (0-0) down (0-12) nc (3-3) down (1-12) nc (9-10)	(DACK)	down (1-13)		up (22-2)
chain down (1-13) IIC (0-0)	M80482, subtilisin-like protein (FACE*)		10-10)	up (17-1)
chain down (0-12) nc (0-0) chain down (0-12) nc (3-3) down (1-12) nc (9-10)	Z26653, laminin M chain (merosin)	down (1-13)		
down (0-12) down (1-12) down (1-12)	**************************************	down (0-12)	nc (0-0)	up (25-0)
down (0-12) nc (9-10)	M30000, 017040, Elastin		nc (3-3)	up (11-0)
down (1-12)	X05610, type IV collagen alpha -2 chain	down (0-12)	(2.2) 211	
Gene	x67325 n27 interferon alpha-inducible gene	down (1-12)	nc (9-10)	up (10-2)

Abbildung

[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name
Vergleich Endometriose versus Normal (sekr. Phase) = Comparison
 of Endometriosis versus Normal (prol. Phase) = Comparison
 of Endometriosis versus Normal (prol. Phase)
Vergleich sekr. versus prol. Phase (Endometrium) = Comparison
 of Secr. versus Prol. Phase (Endometrium)

The state of the s

Datenbank-Nr., Name	Vergleich Endometriose	Vergleich Endometriose	Vergleich sekr. versus prol. Phase
	versus Normal (sekr. Phase)	versus Normal (prol. Phase)	(Endometrium)
D42073, reticulocalbin	down (0-11)	nc (8-5)	up (11-2)
U07919, aldehyde dehydrogenase 6	down (1-11)	nc (13-9)	up (22-0)
U81607, gravin	down (1-11)	nc (8-7)	(18-1)
M30269, nidogen	down (0-10)	nc (8-14)	up (15-3)
D42108, phospholipase C Epsilon	down (1-10)	, nc (12-14)	up (25-0)

Abbildung 1

A,

.

-

THE THE PARTY OF T

Figure 1

[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name

Vergleich Endometriose versus Normal (sekr. Phase) = Comparison

of Endometriosis versus Normal (Secr. Phase)

Vergleich Endometriose versus Normal (prol. Phase) = Comparison

of Endometriosis versus Normal (Prol. Phase)

Vergleich sekr. versus prol. Phase (Endometrium) = Comparison

of Secr. versus Prol. Phase (Endometrium)

The state of the s

(1
	Σ,
Ī	-
7	こここ
7	ב
~	Ċ
_	۰.

DDIIddiig 4		
seq.IDNO	Name	Proteinsequenz
		1
	Fibronektin	
_		YTGNTYRVGD LIERFINDSHI MOTSVAWGET
		AGISIAVGET
		GRITCTSRNR CNDQDTRTSY RIGDTWSKKD NRGNLLQCIC TGNGRGEWRC
		AAVYOPOPHP OPPPYGHCVT DSGVVYSVGM OWLKTOGNKO MLCTCLGNGV
		VI.PFTYNGRT FYSCTTEGRQ DGHLWCSTTS NYEQDQKYSF CTDHTVLVQT
		VTDCTSEGRE
		VIOCIDIO A RECEIDATE
		NGRGEWICIA ISKUNDKOTY
		HGVKIQCICI GRGIGZENICZ
		חווד כל ביו ביו
		VTGETTPFSP LVATSESVTE ITASSFVVSW VSASDTVSGF KVEIELSEEG
		LPGRKYIVNV YQISEDGEQS LILSTSQTTA PDAPPDPTVD QVDDTSIVVR
		SO IN IGNOOD
		GSS16LINDER THUS THUS COMMONDATION
		RAQITGIRDI VGLIMMADA
		EYTVSLVAIK GNQESPKATG VFTTLQPGSS IPPYNTEVTE TTIVITWIFA
		VTSDSGSIVV SGLTPGVEYV YTIQVLRDGQ ERDAPIVNKV VTPLSPPTNL
		PETERSON THE
	•	PDIIGINIII IIIXXX
		TULKETNIGE DIFFICATIONS PTGIDESDIT
		WILL CHARLES
		K K F
, s d		

[Key to Table:]

Proteinsequenz = Protein Sequence Fibronektin = fibronectin

Harry Harry Harry Harry Harry and the state of the state

Seq.IDNO	Name	Proteinsequenz
		PAVTVRYYRI TYGETGGNSP VQEFTVPGSK STATISGLKP GVDYTITVYA VTGRGDSPAS SKPISINYRT
		TDVQDNSISV KWLPSSSPVT GYRVTTTPKN GPGPTKTKTA GPDQTEMTIE
		TISWRTKTET ITGFQVDAVP ANGQTPIQRT IKPDVRSYTI TGLQPGTDYK
		_
		TGOEALSOTT
····		HKVREEVVTV
		CHDNGVNYKI
<u></u>		
		REDSRE
1:	·	
2	Insulin-like	SGGGGGARAE
	growth factor binding protein-2	LAVFREKVTE
		HHLGLEEPKK LRPPPARTPC QQELDQVLER ISTMRLPDER GPLEHLYSLH IPNCDKHGLY NLKQCKMSLN GQRGECWCVN PNTGKLIQGA PTIRGDPECH LFYNEQQEAR GVHTQRMQ

[Key to Table:]

The state of the s

Seq.IDNO	Name	Proteinsequenz
	Transmembrane	MGAARGSPAR PRRLPLLSVL LLPLLGGTQT AIVFIKQPSS QDALQGRRAL LRCEVEAPGP VHVYWLLDGA
7)	recentor PTK7	PVODTERRFA QGSSLSFAAV DPLQDSGTFQ CVARDDVTGE EARSANASFN IKWIEAGPVV LKHPASEAEI
		SSONFTLSIA DESFARVVLA PQDVVVARYE EAMFHCQFSA QPPPSLQWLF EDETPITNRS RPPHLRRATV
		FANGSLLLTO VRPRNAGIYR CIGOGORGPP IILEATLHLA EIEDMPLFEP RVFTAGSEER VTCLPPKGLP
		EPSVWWEHAG VRLPTHGRVY QKGHELVLAN IAESDAGVYT CHAANLAGOR RQDVNITVAT VPSWLKKPQD
		SQLEEGKPGY LDCLTQATPK PTVVWYRNQM LISEDSRFEV FKNGTLRINS VEVYDGTWYR CMSSTPAGSI
		EAQAVLQVLE KLKFTPPPQP QQCMGFDKEA TVPCSATGRE KPTIKWERAD GSSLPEWVTD NAGTLHFARV
		TRDDAGNYTC IASNGPQGQI RAHVQLTVAV FITFKVEPER TTVYQGHTAL LQCEAQGDPK PLIQWKGKDR
		SLGSGPAATN KRHSTSDKMH FPRSSLQPIT TLGKSEFGEV FLAKAQGLEE GVAETLVLVK SLQSKDEQQQ
		LDFRRELEMF GKLNHANVVR LLGLCREAEP HYMVLEYVDL EDLKQFLRIS KSKDEKLKSQ PLSTKQKVAL
		CTOVALGMEH LSNNRFVHKD LAARNCLVSA QRQVKVSALG LSKDVYNSEY YHFRQAWVAL RWMSPEAILE
		GDFSTKSDVW ASGVLMWEVF THGEMPHGGQ ADDEVLADLQ AGKARLPQPE GCPSKLYRLM QRCWALSPKD
		RPSFSEIASA LGDSTVDSKP
	Platelet-derived	MGTSHPAFLV LGCLLTGLSL ILCQLSLPSI LPNENEKVVQ LNSSFSLRCF GESEVSWQYP MSEEESSDVE
1 1	growth factor	IRNEENNSGL FVTVLEVSSA SAAHŢGLYTC YYNHTQTEEN ELEGRHIYIY VPDPDVAFVP LGMTDYLVIV
	or alpha	EDDDSAIIPC RTTDPETPVT LHNSEGVVPA SYDSRQGFNG TFTVGPYICE ATVKGKKFQT IPFNVYALKA

The state of the s

[Key to Table:]

The first prince that the first prince from the first prince prin

Seq.IDNO	Name	Proteinsequenz
		IMATIX ADMINIST MANAGEMENT
******		TVKDSGDYEC AARQATREVK EMKKVTISVH EKGFIEIKPT FSQLEAVNLH EVKHFVVEVR AYPPPRISML
		VODHHGSTGG OTVRCTAEGT PLPDIEWMIC KDIKKCNNET SWTILANNVS NIITEIHSRD RSTVEGRVTF
		NPADESTRSY
		LISDDNSEGL TLLDLLSFTY
		SKGSTFLPVK
		VDSDNAYIG VTYKNE
		SSQTSEESAI ETGSSSSTFI KREDETIEDI DMMDDIGIDS SDLVEDSFL
,	Collagen type	GEVGADGIPG FPGLPGREGI AGPQGPKGDR GSRGEKGDPG KDGLGQPGLP GPKGPFGFVV IVSEKDGSVI
٠	ת מ	SVPGPEGRRG FAGFPGPAGP KGNLGSKGEL GSPGPKGEKG EPGSIFSPDG GALGPAQKGA KGEPGFRGPP
	• .	
		1

[Key to Table:]

The time of the first of the fi

. Hilly . J. Carly . 1987		
	Proteinsequenz	
	Name	
Abbildung 2	Seq.IDNO	

		IVRRADRAAV PIVNLKUELL FPSWEALFSG SEGFENFGAR IFSTEGIEVE IN THE STOP	
		LTESYCETWR TEAPSATGQA SSLLGGRLLG QSAASCHHAY IVLCIENSFM TASK	K
	Subtilisin-like	MPPRAPPAPG PRPPPRAAAA TDTAAGAGGA GGAGGAGGPG FRPLAPRPWR WLI	WLLLLALPAA CSAPPRRVY
	nrotein (PACE4	GPAEADRVAA AHGYLNLGQI GNLEDYYHFY HSKTFKRSTL	SSRGPHTFLR MDPQVKWLQQ
		VRSDPQALYF NDPIWSNMWY LHCGDKNSRC RSEMNVQAAW	KRGYTGKNVV VTILDDGIER
		SYASYDVNGN DYDPSPRYDA SNENKHGTRC AGEVAASANN	SYCIVGIAYN AKIGGIRMLD
		SLGIRPNYID IYSASWGPDD DGKTVDGPGR LAKQAFEYGI	KKGRQGLGSI FVWASGNGGR
		EGDYCSCDGY TNSIYTISVS SATENGYKPW YLEECASTLA TTYSSGAFYE RK	RKIVTTDLRQ RCTDGHTGTS
		VSAPMVAGII ALALEANSQL TWRDVQHLLV KTSRPAHLKA SDWKVNGAGH KV	KVSHFYGFGL VDAEALVVEA
*****		KKWTAVPSQH MCVAASDKRP RSIPLVQVLR TTALTSACAE HSDQRVVYLE HV	HVVVRTSISH PRRGDLQIYL
		LAKRLLDLSN EGFTNWEFMT VHCWGEKAEG QWTLEIQDLP	SQVRNPEKQG KLKEWSLILY
		GTAEHPYHTF SAHQSRSRML ELSAPELEPP KAALSPSQVE VPEDEEDYTA QS	QSTPGSANIL QTSVCHPECG
		DKGCDGPNAD QCLNCVHFSL GSVKTSRKCV SVCPLGYFGD TAARRCRRCH KG	KGCETCSSRA ATQCLSCRRG
		FYHHQEMNTC VTLCPAGFYA DESQKNCLKC HPSCKKCVDE PEKCTVCKEG FS	FSLARGSCIP DCEPGTYFDS
		TCGTCVGPGR EECIHCAKNF HFHDWKCVPA CGEGFYPEEM	PGLPHKVCRR CDENCLSCAG
		SSRNCSRCKT GFTQLGTSCI TNHTCSNADE TFCEMVKSNR LCERKLFIQF CC	CCRTCLLAG
7	Laminin M chain	MPGAAGVLLL LLLSGGLGGV QAQRPQQQRQ SQAHQQRGLF PAVLNLASNA	LITTNATCGE KGPEMYCKLV
~	ı d	EHVPGQPVRN PQCRICNQNS SNPNQRHPIT NAIDGKNTWW QSPSIKNGIE YH	YHYVTITLDL QQVFQIAYVI
		NWILERSLDD VEYKPWQYHA VTDTECLTLY NIYPRTGPPS	YAKDDEVICT SFYSKIHPLE
		GRPSADDPSP ELLEFTSARY IRLRFQRIRT LNADLMMFAH	KDPREIDPIV TRRYYYSVKD
		ISVGGMCICY GHARACPLDP ATNKSRCECE HNTCGDSCDQ CCPGFHQKPW RA	RAGTFLTKTE CEACNCHGKA

[Key to Table:]

The first that the fi

Ç	7
ζ	7
2	
	\supset
3	\supseteq
Ξ	_
ک	\Box
ک	\supseteq
<	1
	•

CTQNTAGINC ETCTDGFFRP K VSCDRCARGY TGYPDCKACN C SGVSNRCQSS YWTYGKIQDM S LGNKLPAVGG QLTFTISYDL E TIHGTHFPVR RKEFWTVLAN L TGSSCESCWP RHRRVNGTIF G SEDCQPCACP LNIPSNNFSP T LDFSIPGSCD SLSGSCLICK F LDFSIPGSCD CKAGTFGLQS P ANVQGQRCDK CKAGTFGLQS P GKFGLDAKNP LGCSSCYCFG T MDLMREDLHL EPFYWKLPEQ F MDLMREDLHL EPFYWKLPEQ F WAAAPLIGQLT RHEIEMTEKE F VAEQGRGTTM TPPADLIEKC F PETSICQNCQ HHTAGDFCER G PETSICQNCQ HHTAGDFCER G VAEQGRGTTM TPPADLIEKC F NTLVTEMNEL LTRATKVTAD ERNLEGLQKE IDQMIKELRR NKVDDAMDLL REATDKIREA			
RRNLSIANIRG KYIGGGVCIN CTQNTAGINC ETCTDGFFRP K KHARRGLAPG SCHCKTGFGG VSCDRCARGY TGYPDCKACN C KSGFFNLQED NWKGCDECFC SGVSNRCQSS YWTYGKIQDM S NAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL B VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEPMTVLAN L YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF G GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP T YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F VYGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P VTGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY F TGQCTCKVNV GGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQCTCKVNV GGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC F GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER F SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP F SPSCVAEGLD DYRCTACPRG YEGGYCERCA PGYTGSPGNP F SPSCVAEGLD DYRCTACPRG YEGGY T SPSCVAEGLD T SPSCVAEGLD T SPSCVAEGLD T SPGNAEGLD T SPGN	Seq.IDNO	Name	Proteinsequenz
RRNLSIANIRG KYIGGGVCIN CTQNTAGINC ETCTDGFFRP K KHARRGLAPG SCHCKTGFGG VSCDRCARGY TGYPDCKACW C KSGFFNLQED NWKGCDECFC SGVSNRCQSS YWTYGKIQDM S NAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL B VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFMTVLAN L YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRWMGTIF G GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP T YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F VYGGPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS F VTGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY F TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQCTCKVNV GGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQCTCKVNV GGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQCTCKVNV GGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQTTCCVVQ QCNGHSSLCD PETSICQNCQ HHTAGDFCER F SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP F SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP F SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP F SPSCVAEGLID DYRCTACPRG YEGGYCERCA PGYTGSPGNP F SPSCVAEGLID DYRCTACPRG YEGYCERCA PATDKIRGNP F SPSCVAEGLID DYRCTACPRO YEGGYCERCA PGYTGSPGNP			
KHARRGLAPG SCHCKTGFGG VSCDRCARGY TGYPDCKACN C KSGFFNLQED NWKGCDECFC SGVSNRCQSS YWTYGKIQDM S NAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL B VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFMTVLAN L YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRWGTIF G GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP T GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP T YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P TYGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY E TGQTTCKNNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T RALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYMKLPEQ I RALGHTTTKGI VFQHPEIVAH MDLMREDLHL TPPADLIEKC E GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER E SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP E GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE E R ANNEKAIKLN ETLGILAEGNL NTLVTEMNEL LTRATKVTAD E B ANNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR E SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA	,		RRNLSLNIRG KYIGGGVCIN CTQNTAGINC
KSGFFNLQED NWKGCDECFC SGVSNRCQSS YWTYGKIQDM SNAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL B VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFWTVLAN LYPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TGEPYCCLP GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TYFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK FORGKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC INCOPKTGRC INCOPKTGRC GYTGGCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG GYTGGTTTKGI VFQHPEIVAH MDLMREDLHL EPFYMKLPEQ GYTGGTTTKGI VFQHPEIVAH MAAPLIGQLT RHEIEMTEKE GYNQUIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE GYNGHSSLCD PFTSICQNCQ HHTAGDFCER GYTGTTCVPC QCNGHSSLCD PFTSICQNCQ HHTAGDFCER GYTGTTCVPC QCNGHSSLCD PFTSICQNCQ HHTAGDFCER GYTGTTCVPC CHWHAREGWE CVFCGDECTG LLLGDLARLE GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE GYNDEARLE LTRATKVTAD ESRGENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			KHARRGLAPG SCHCKTGFGG VSCDRCARGY
NAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL B VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFMTVLAN L YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF G GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP T YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P TGQCTCKVNV GGLHCDRCRP GKFGLDAKNP LGCSSCYCFG T TGQCTCKVNV GGHCDRCRP GKFGLDAKNP LGCSSCYCFG T NPQVIIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE F INATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC F GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER F SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP F GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE F K HLLSPQRAPE RLIQLAEGNL NTLYTEMNEL LTRATKVTAD E AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR E SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			KSGFFNLQED NWKGCDECFC SGVSNRCQSS
YYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFMTVLAN LYPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TYFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK BNAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS PYGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQTTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ I IKATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC GGTTGTTGOTC HHTAGDFCER GATGRCCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE GATGRCCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE ANNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			NAEARQALPH SYYWSAPAPY LGNKLPAVGG
YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TYFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK FORGKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC INCKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG GPTLGTTKGI VFQHPEIVAH MAAPLIGQLT RHEIEMTEKE GPTLGTTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE ANNBEAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SROBENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			VYLHPSEEHT NVLLLKEESF TIHGTHFPVR
GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TYFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK BNAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS PVGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TYGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TYGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG GALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYMKLPEQ GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE GHLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD SROWNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			YPTDGSIAAA VEVCQCPPGY TGSSCESCWP
YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK F NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P VTGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY E TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T ALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ I I NPQVIIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE V GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER G SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP G GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE K HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD S AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR E SRGENEEMEK DIREKLADYK NKVDDAWDLL REATDKIREA			GGPYCDKCLP GFYGEPTKGT SEDCQPCACP
NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS P VTGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY F TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG T ALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ I IKATYGNFMR QSRISBISME VAEQGRGTTM TPPADLIEKC I GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER G SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP G CHLLSPQRAPE KLIQLAEGNL NTLVTEMNEL LTRATKVTAD THALSPQRAPE KLIQLAEGNL NTLVTEMNEL LTRATKVTAD SRGENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD
VTGKKCDRCA HGYFNFQEGG CTACECSHLG NNCDPKTGRC I NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY FGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ I NPQVIIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD ANNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			NAGGSFSEVC HSQTGQCECR ANVQGQRCDK
TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ INPQVIIRGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD SCHEENER DIREKLADYK NKVDDAMDLL REATDKIREA			VTGKKCDRCA HGYFNFQEGG CTACECSHLG
TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG ALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ INPQVIIRGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE INFATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC INCATYGNFMC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD ANNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC
ALQHTTTKGI VFQHPEIVAH MDLMREDLHL EPFYWKLPEQ INPQVIIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE IKATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC IGPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP
NPQVIIRGGT PTHARIIVRH MAAPLIGQLT RHEIEMTEKE I IKATYGNFMR QSRISBISME VAEQGRGTTM TPPADLIEKC I GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE CHLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			ALQHTTTKGI VFQHPEIVAH MDLMREDLHL
IKATYGNFMR QSRISEISME VAEQGRGTTM TPPADLIEKC I GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER I SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP I GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE ILLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD I AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR ISRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			NPQVIIRGGT PTHARIIVRH MAAPLIGQLT
GPTLGTCVPC QCNGHSSLCD PETSICQNCQ HHTAGDFCER SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			IKATYGNFMR QSRISEISME VAEQGRGTTM
SPSCVAEGLD DYRCTACPRG YEGQYCERCA PGYTGSPGNP GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			GPTLGTCVPC QCNGHSSLCD PETSICQNCQ
GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			SPSCVAEGLD DYRCTACPRG YEGQYCERCA
HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			GATGRKCDGC KHWHAREGWE CVFCGDECTG
AVNEKAIKLN ETLGTRDEAF ERNLEGLOKE IDQMIKELRR SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA			HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL
SRGENEEMEK DLREKLADYK NKVDDAMDLL REATDKIREA			AVNEKAIKLN ETLGTRDEAF ERNLEGLOKE
			SRGENEEMEK DLREKLADYK NKVDDAWDLL
SIIDYVEDIQ TKLPPMSEEL			

The state of the s

[Key to Table:]

Seq.IDNO	Name	Proteinsequenz
		SQAESHAAQL NDSSAVLDGI LDEAKNISFN ATAAFKAYSN IKDYIDEAEK VAKEAKDLAH EATKLATGPR
		GLLKEDAKGC LQKSFRILNE AKKLANDVKE NEDHLNGLKT RIENADARNG DLLRTLNDTL GKLSAIPNDT
		AAKLQAVKDK ARQANDTAKD VLAQITELHQ NLDGLKKNYN KLADSVAKTN AVVKDPSKNK IIADADATVK
		NLEQEADRLI DKLKPIKELE DNLKKNISEI KELINQARKQ ANSIKVSVSS GGDCIRTYKP EIKKGSYNNI
		VVNVKTAVAD NLLFYLGSAK FIDFLAIEMR KGKVSFLWDV GSGVGRVEYP DLTIDDSYWY RIVASRTGRN
		GTISVRALDG PKASIVPSTH HSTSPPGYTI LDVDANAMLF VGGLTGKLKK ADAVRVITFT GCMGETYFDN
		KPIGLWNFRE KEGDCKGCTV SPQVEDSEGT ATRDLRDFMS VELTDGHIKV SYDLGSGMAS VVSNQNHNDG
		KWKSFTLSRI QKQANISIVD IDTNQEENIA TSSSGNNFGL DLKADDKIYF GGLPTLRNLS MKARPEVNLK
<u> </u>		KYSGCLKDIE ISRTPYNILS SPDYVGVTKG CSLENVYTVS FPKPGFVELS PVPIDVGTEI NLSFSTKNES
		GIILLGSGGT PAPPRRKRRQ TGQAYYVILL NRGRLEVHLS TGARTMRKIV IRPEPNLFHD GREHSVHVER
		TRGIFTVQVD ENRRYMQNLT VEQPIEVKKL FVGGAPPEFQ PSPLRNIPPF EGCIWNLVIN SVPMDFARPV
		SFKNADIGRC AHOKLREDED GAAPAEIVIQ PEPVPTPAFP TPTPVLTHGP CAAESEPALL IGSKQFGLSR
		NSHIAIAFDD TKVKNRLTIE LEVRTEAESG LLFYMAAINH ADFATVQLRN GLPYFSYDLG SGDTHTMIPT
		KINDGQWHKI KIMRSKQEGI LYVDGASNRT ISPKKADILD VVGMLYVGGL PINYTTRRIG PVTYSIDGCV
		RNLHMAEAPA DLEQPTSSFH VGTCFANAQR GTYFDGTGFA KAVGGFKVGL DLLVEFEFAT TTTTGVLLGI
		SSOKMDGMGI EMIDEKLMFH VDNGAGRFTA VYDAGVPGHL CDGQWHKVTA NKIKHRIELT VDGNQVEAQS
		PNPASTSADT NDPVFVGGFP DDLKQFGLTT SIPFRGCIRS LKLTKGTASH WRLILPRPWN
8	Elastin	MAGLTAAAPR PGVLLLLLSI LHPSRPGGVP GAIPGGVPGG VFYPGAGLGA LGGGALGPGG KPLKPVPGGL
		AGAGLGAGLG AFPAVTFPGA LVPGGVADAA AAYKAAKAGA GLGGVPGVGG LGVSAGAVVP QPGAGVKPGK
<i>-</i>		VPGVGLPGVY PGGVLPGARF PGVGVLPGVP TGAGVKPKAP GVGGAFAGIP GVGPFGGPQP GVPLGYPIKA
		PKLPGGYGLP YTTGKLPYGY GPGGVAGAAG KAGYPTGTGV GPQAAAAAA KAAAKFGAGA AGVLPGVGGA

[Key to Table:]

The transport of the state of t

`	٦
ζ	J
	_
_	J
τ	7
=	=
۷	\supset
2	\supset
<	

Seq.IDNO	Name	·	Proteinsequenz	aquenz					
			GVPGVPGAIP	GIGGIAGVGT	PAAAAAAAA	AKAAKYGAAA	GLVPGGPGFG	PGVVGVPGAG	VPGVGVPGAG
			IPVVPGAGIP	GAAVPGVVSP	Eaaakaaaka	AKYGARPGVG	VGGIPTYGVG	AGGFPGFGVG	VGGIPGVAGV
			PSVGGVPGVG	GVPGVGISPE	aqaaaakaa	KYGVGTPAAA	aakaaakaaq	FALLINLAGLV	PGVGVAPGVG
·			VAPGVGVAPG	VGLAPGVGVA	PGVGVAPGVG	VAPGIGPGGV	AAAAKSAAKV	AAKAQLRAAA	GLGAGIPGLG
	***************************************		VGVGVPGLGV	GAGVPGLGVG	AGVPGFGAVP	GALAAAKAAK	YGAAVPGVLG	GLGALGGVGI	PGGVVGAGPA
	14 day 2		AAAAAAKAAA	KAAQFGLVGA	AGLGGLGVGG	LGVPGVGGLG	GIPPAAAAKA	AKYGAAGLGG	VLGGAGQFPL
			GGVAARPGFG	LSPIFPGGAC	LGKACGRKRK				
6	Alpha-2 type	pe IV	MGRDQRAVAG	PALRRWLLLG	TVTVGFLAQS	VLAGVKKFDV	PCGGRDCSGG	CQCYPEKGGR	GQPGPVGPQG
	collagen		YNGPPGLQGF	PGLQGRKGDK	GERGAPGVTG	PKGDVGARGV	SGFPGADGIP	GHPGQGGPRG	RPGYDGCNGT
			QGDSGPQGPP	GSEGFTGPPG	PQGPKGQKGE	PYALPKEERD	RYRGEPGEPG	LVGFQGPPGR	PGHVGQMGPV
			GAPGRPGPPG	PPGPKGQQGN	RGLGFYGVKG	EKGDVGQPGP	NGIPSDTLHP	IIAPTGVTFH	PDQYKGEKGS
			EGEPGIRGIS	LKGEEGIMGF	PGLRGYPGLS	GEKGSPGQKG	SRGLDGYQGP	DGPRGPKGEA	GDPGPPGLPA
-			YSPHPSLAKG	ARGDPGFPGA	QGEPGSQGEP	GDPGLPGPPG	LSIGDGDQRR	GLPGEMGPKG	FIGDPGIPAL
			YGGPPGPDGK	RGPPGPPGLP	GPPGPDGFLF	GLKGAKGRAG	FPGLPGSPGA	RGPKGWKGDA	GECRCTEGDE
		•	AIKGLPGLPG	PKGFAGINGE	PGRKGDKGDP	GOHGLPGFPG	LKGVPGNIGA	PGPKGAKGDS	RTITTKGERG
			QPGVPGVPGM	KGDDGSPGRD	GLDGFPGLPG	PPGDGIKGPP	GDPGYPGIPG	TKGTPGEMGP	PGLGLPGLKG
	-4		QRGFPGDAGL	PGPPGFLGPP	GPAGTPGQID	CDTDVKRAVG	GDRQEAIQPG	CIAGPKGLPG	LPGPPGPTGA
			KGLRGIPGFA	GADGGPGPRG	LPGDAGREGF	PGPPGFIGPR	GSKGAVGLPG	PDGSPGPIGL	PGPDGPPGER
			GLPGEVLGAQ	PGPRGDAGVP	GQPGLKGLPG	DRGPPGFRGS	QGMPGMPGLK	GQPGLPGPSG	QPGLYGPPGL
			HGFPGAPGQE	GPLGLPGIPG	REGLPGDRGD	PGDTGAPGPV	GMKGLSGDRG	DAGFTGEQGH	PGSPGFKGID
			GMPGTPGLKG	DRGSPGMDGF	QGMPGLKGRP	GFPGSKGEAG	FFGIPGLKGL	AGEPGFKGSR	GDPGPPGPPP

[Key to Table:]

Seq.IDNO	Name	Proteinsequenz
		WILDGMKDIK GEKGDEGPMG LKGYLGAKGI QGMPGIPGLS GIPGLPGRPG HIKGVKGDIG VPGIPGLPGF
		GFPGFIGSRG
		FGEKGTEGDI GFPGITGVTG VQGPPGLKGQ TGFPGLTGPP GSQGELGRIG LPGGKGDDGW PGAPGLPGFP
		GLRGIRGLHG LPGTKGFPGS PGSDIHGDPG FPGPPGERGD PGEANTLPGP VGVPGQKGDQ GAPGERGPPG
		SPGLQGFPGI TPPSNISGAP GDKGAPGIFG LKGYRGPPGP PGSAALPGSK GDTGNPGAPG TPGTKGWAGD
		SGPQGRPGVF GLPGEKGPRG EQGFMGNTGP TGAVGDRGPK GPKGDPGFPG APGTVGAPGI AGIPQKIAIQ
		PGTVGPQGRR GPPGAPGEIG PQGPPGEPGF RGAPGKAGPQ GRGGVSAVPG FRGDEGPIGH QGPIGQEGAP
		GRPGSPGLPG MPGRSVSIGY LLVKHSQTDQ EPMCPVGMNK LWSGYSLLYF EGQEKAHNQD LGLAGSCLAR
		FSTMPFLYCN PGDVCYYASR NDKSYWLSTT APLPMMPVAE DEIKPYISRC SVCEAPAIAI AVHSQDVSIP
		HCPAGWRSLW IGYSFLMHTA AGDEGGGQSL VSPGSCLEDF RATPFIECNG GRGTCHYYAN KYSFWLTTIP
		EQSFQGSPSA DTLKAGLIRT HISRCQVCMK NL
1.0	p27	MEASALTSSA VTSVAKVVRV ASGSAVVLPL ARIATVVIGG VVAMAAVPMV LSAMGFTAAG IASSSIAAKM
) {		MSAAAIANGG GVASGSLVGT LQSLGATGLS GLTKFILGSI GSAIAAVIAR FY
-	Reticulocalbin	MARGGRGRRL GLALGLLLAL VLAPRVLRAK PTVRKERVVR PDSELGERPP EDNQSFQYDH EAFLGKEDSK
+ +		TFDQLTPDES KERLGKIVDR IDNDGDGFVT TEELKTWIKR VQKRYIFDNV AKVWKDYDRD KDDKISWEEY
		KQATYGYYLG NPAEFHDSSD HHTFKKMLPR DERRFKAADL NGDLTATREE FTAFLHPEEF EHMKEIVVLE
	•	TLEDIDKNGD GFVDQDEYIA DMFSHEENGP EPDWVLSERE QFNEFRDLNK DGKLDKDEIR HWILPQDYDH
		AQAEARHLVY ESDKNKDEKL TKEEILENWN MFVGSQATNY GEDLTKNHDE L
(bldehvde	MATANGAVEN GOPDGKPPAL PRPIRNLEVK FTKIFINNEW HESKSGKKFA TCNPSTREQI CEVEEGDKPD
7	dehydrogenase 6	
	•	

[Key to Table:]

Seq.IDNO	Name	Proteinsequenz
		CHULL ROOMER'S CHILDREN
		LTALYLGSLI KEAGFPPGVV NIVPGFGPTV GAAISSHPQI NKIAFTGSTE VGKLVKEAAS RSNLKRVTLE
		LGGKNPCIVC ADADLDLAVE CAHQGVFFNQ GQCCTAASRV FVEEQVYSEF VRRSVEYAKK RPVGDPFDVK
		TEQGPQIDQK QFDKILELIE SGKKEGAKLE CGGSAMEDKG LFIKPTVFSE VTDNMRIAKE EIFGPVQPIL
		KFKSIEEVIK RANSTDYGLT AAVFTKNLDK ALKLASALES GTVWINCYNA LYAQAPFGGF KMSGNGRELG
		EYALAEYTEV KTVTIKLGDK NP
1.2	Gravin	MGAGSSTEQR SPEQPPEGSS TPAEPEPSGG GPSAEAAPDT TADPAIAASD PATKLLQKNG QLSTINGVAE
		QDELSLQEGD LNGQKGALNG QGALNSQEEE EVIVTEVGQR DSEDVSERDS DKEMATKSAV VHDITDDGQE
		ENRNIEQIPS SESNLEELTQ PTESQANDIG FKKVFKFVGF KFTVKKDKTE KPDTVQLLTV KKDEGEGAAG
		AGDHODPSLG AGEAASKESE PKOSTEKPEE TLKREQSHAE ISPPAESGQA VEECKEEGEE KQEKEPSKSA
		ESPTSPVTSE TGSTFKKFFT QGWAGWRKKT SFRKPKEDEV EASEKKKEQE PEKVDTEEDG KAEVASEKLT
		ASEQAHPQEP AESAHEPRLS AEYEKVELPS EEQVSGSQGP SEEKPAPLAT EVFDEKIEVH QEEVVAEVHV
	· · · · · · · · · · · · · · · · · · ·	STVEERTEEQ KTEVEETAGS VPAEELVGMD AEPQEAEPAK ELVKLKETCV SGEDPTQGAD LSPDEKVLSK
		PPEGVVSEVE MLSSQERMKV QGSPLKKLFT STGLKKLSGK KQKGKRGGGD EESGEHTQVP ADSPDSQEEQ
		KGESSASSPE EPEEITCLEK GLAEVQQDGE AEEGATSDGE KKREGVTPWA SFKKMVTPKK RVRRPSESDK
		EDELDKVKSA TLSSTESTAS EMQEEMKGSV EEPKPEEPKR KVDTSVSWEA LICVGSSKKR ARRRSSSDEE
		GGPKAMGGDH QKADEAGKDK ETGTDGILAG SQEHDPGQGS SSPEQAGSPT EGEGVSTWES FKRLVTPRKK
	······································	SKSKLEEKSE DSIAGSGVEH STPDTEPGKE ESWVSIKKFI PGRRKKRPDG KQEQAPVEDA GPTGANEDDS
		DVPAVVPLSE YDAVEREKME AQQAQKGAEQ PEQKAATEVS KELSESQVHM MAAAVADGTR AATIIEERSP
		SWISASVTEP LEQVEAEAAL LTEEVLEREV IAEEEPPTVT EPLPENREAR GDTVVSEAEL TPEAVTAAET
		AGPLGSEEGT EASAAEETTE MVSAVSQLTD SPDTTEEATP VQEVEGGVPD IEEQERRTQE VLQAVAEKVK
,3		משטשטשטשט אין

[Key to Table:]

STAIN SEAST SEAST SERVICE OF THE SEASON SEAS

•	*
ζ	J.
0	_
-	=
7	J
=	☱
2	⋽
_	$\bar{\beth}$
~	7
	•

EESQLPGTGG PEDVLQ KAPQVTESIE SSELVT VEIHEENEVA SGTQSG EGTQEADQYA DEKTKD PVEREMVVQV EREKTE QEEAVCTKIQ VQSSEA AVPTGPDCQA KSTPVI LETKSSKLVQ NIIQTP TSAKEESEST AVGQAI ALLAERIEKS LVEPKE HSESDKAITP QAQEEI IDAVYVTTNG IIATSI GFPEISFQPS SAVVV ENNQVPAVVA FSQGS LGTEDGAEYD DEDED	Proteinsequenz
RAPQVTESIE S VEIHEENEVA S EGTQEADQYA D PVEREMVVQV E GEAVCTKIQ V AVPTGPDCQA P LETKSSKLVQ P TSAKEESEST P ALLAERIEKS I HSESDKAITP (IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	- 1
KAPQVTESIE S VEIHEENEVA S EGTQEADQYA E PVEREMVVQV E QEBAVCTKIQ V AVPTGPDCQA H LETKSSKLVQ N TSAKEESEST J ALLAERIEKS I HSESDKAITP G IDAVYVTTNG GFPEISFQPS ENNQVPAVVA	EESQLPGTGG PEDVLQPVQR AEAERPEEQA EASGLKKETD VVLKVDAQEA KTEPFTQGKV VGQTTPESFE
VETHEENEVA S EGTQEADQYA E PVEREMVVQV E QEEAVCTKIQ V AVPTGPDCQA H LETKSSKLVQ P TSAKEESEST P ALLAERIEKS I HSESDKAITP (IDAVYVTTNG GFPEISFQPS ENNQVPAVVA	KAPQVTESIE SSELVTTCQA ETLAGVKSQE MVMEQAIPPD SVETPTDSET DGSTPVADFD APGTTQKDEI
EGTQEADQYA E PVEREMVVQV E QEEAVCTKIQ V AVPTGPDCQA P LETKSSKLVQ N TSAKEESEST P ALLAERIEKS I HSESDKAITP G GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	VEIHEENEVA SGTQSGGTEA EAVPAQKERP PAPSSFVFQE ETKEQSKMED TLEHTDKEVS VETVSILSKT
PVEREMVVQV EQEEAVCTKIQ V AVPTGPDCQA H LETKSSKLVQ N TSAKEESEST J ALLAERIEKS I HSESDKAITP (HSESDKAITP (IDAVYVTTNG GFPEISFQPS ENNQVPAVVA	EGTQEADQYA DEKTKDVPFF EGLEGSIDTG ITVSREKVTE VALKGEGTEE AECKKDDALE LQSHAKSPPS
QEEAVCTKIQ V AVPTGPDCQA H LETKSSKLVQ P TSAKEESEST P ALLAERIEKS I HSESDKAITP (HSESDKAITP (GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	PVEREMVVQV EREKTEAEPT HVNEEKLEHE TAVTVSEEVS KQLLQTVNVP IIDGAKEVSS LEGSPPPCLG
AVPTGPDCQA F LETKSSKLVQ N TSAKEESEST N ALLAERIEKS I HSESDKAITP C HSESDKAITP C IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	QEEAVCTKIQ VQSSEASFTL TAAAEEEKVL GETANILETG ETLEPAGAHL VLEEKSSEKN EDFAAHPGED
TSAKEESEST 7 ALLAERIEKS 1 HSESDKAITP C HSESDKAITP C IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	AVPTGPDCQA KSTPVIVSAT TKKGLSSDLE GEKTTSLKWK SDEVDEQVAC QEVKVSVAIE DLEPENGILE
TSAKEESEST ALLAERIEKS I HSESDKAITP (HSESDKAITP (IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	LETKSSKLVQ NIIQTAVDQF VRTEETATEM LTSELQTQAH VIKADSQDAG QETEKEGEEP QASAQDETPI
4 MLASSSRIRA IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	TSAKEESEST AVGQAHSDIS KDMSEASEKT MTVEVEGSTV NDQQLEEVVL PSEEEGGGAG TKSVPEDDGH
HSESDKAITP (MLASSSRIRA IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	ALLAERIEKS LVEPKEDEKG DDVDDPENQN SALADTDASG GLTKESPDTN GPKQKEKEDA QEVELQEGKV
MLASSSRIRA I DAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	HSESDKAITP QAQEELQKQE RESAKSELTE S
4 MLASSSRIRA IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	
4 MLASSSRIRA IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	
IDAVYVTTNG GFPEISFQPS ENNQVPAVVA LGTEDGAEYD	MLASSSRIRA AWTRALLLPL LLAGPVGCLS RQELFPFGPG QGDLELEDGD DFVSPALELS GALRFYDRSD
	IDAVYVTTNG IIATSEPPAK ESHPGLFPPT FGAVAPFLAD LDTTDGLGKV YYREDLSPSI TQRAAECVHR
	GFPEISFQPS SAVVVTWESV APYQGPSRDP DQKGKRNTFQ AVLASSDSSS YAIFLYPEDG LQFHTTFSKK
	ENNQVPAVVA FSQGSVGFLW KSNGAYNIFA NDRESIENLA KSSNSGQQGV WVFEIGSPAT TNGVVPADVI
RSFOLAVETF HOOHP	RSFQLAVETF HQQHPQVIDV DEVEETGVVF SYNTDSRQTC ANNRHQCSVH AECRDYATGF CCSCVAGYTG

He died from the first the

[Key to Table:]

The state of the s

•	4
(J.
ς	_
	\exists
٦	J
	=
ک۔	\Box
ک	\supseteq
<	Q

Seq.IDNO	Name	Proteinsequenz
		1
		QQLSVDSVFV LYNQEEKILR YAFSNSIGPV REGSPDALQN PCYIGTHGCD TNAACRPGPR TQFTCECSIG
		FRGDGRTCYD IDECSEQPSV CGSHTICNNH PGTFRCECVE GYQFSDEGTC VAVVDQRPIN YCETGLHNCD
		IPQRAQCIYT GGSSYTCSCL PGFSGDGQAC QDVDECQPSR CHPDAFCYNT PGSFTCQCKP GYQGDGFRCV
		PGEVEKTRCQ HEREHILGAA GATDPQRPIP PGLFVPECDA HGHYAPTQCH GSTGYCWCVD RDGREVEGTR
		TRPGMTPPCL STVAPPIHQG PAVPTAVIPL PPGTHLLFAQ TGKIERLPLE GNTMRKTEAK AFLHVPAKVI
		IGLAFDCVDK MVYWTDITEP SIGRASLHGG EPTTIIRQDL GSPEGIAVDH LGRNIFWTDS NLDRIEVAKL
		DGTQRRVLFE TDLVNPRGIV TDSVRGNLYW TDWNRDNPKI ETSYMDGTNR RILVQDDLGL PNGLHFDAFS
		SQLCWVDAGT NRAECLNPSQ PSRRKALEGL QYPFAVTSYG KNLYFTDWKM NSVVALDLAI SKETDAFQPH
		KOTRLYGITT ALSQCPQGHN YCSVNNGGCT HLCLATPGSR TCRCPDNTLG VDCIERK
۲	Phospholipase C	MPSEKKISSA NDCISFMQAG CELKKVRPNS RIYNRFFTLD TDLQALRWEP SKKDLEKAKL DISAIKEIRL
٦ - ا	Epsilon	GKNTETFTNN GLADQICEDC AFSILHGENY ESLDLVANSA DVANIWVSGL RYLVSRSKQP LDFMEGNQNT
		PRFMWLKTVF EAADVDGNGI MLEDTSVELI KQLNPTLKEA KIRLKFKEIQ KSKEKLTTRV TEEEFCEAFC
		ELCTRPEVYF LLVQISKNKE YLDANDLMLF LEAEQGVTHI TEDICLDIIR RYELSEEGRQ KGFLAIDGFT
		QYLLSSECDI FDPEQKKVAQ DMTQPLSHYY INASHNTYLI EDQFRGPADI NGYIRALKMG CRSVELDVSD
		GSDNEPILCN RNNMTTHVSF RSVIEVINKF AFVASEYPLI LCLGNHCSLP QQKVMAQQMK KVFGNKLYTE
		APLPSESYLP SPEKLKRMII VKGKKLPSDP DVLEGEVTDE DEEAQMSRRM SVDYNGEQKQ IRLCRELSDL
		VSICKSVQYR DFELSMKSQN YWEMCSFSET EASRIANEYP EDFVNYNKKF LSRIYPSAMR IDSSNLNPQD
		FWNCGCQIVA MNFQTPGPMM DLHTGWFLQN GGCGYVLRPS IMRDEVSYFS ANTKGILPGV SPLALHIKII
	,	SGONFPKPKG ACAKGDVIDP YVCIEIHGIP ADCSEQRTKT VQQNSDNPIF DETFEFQVNL PELAMIRFVV
		LDDDYIGDEF IGQYTIPFEC LQPGYRHVPL RSFVGDIMEH VTLFVHIAIT NRSGGGKAQK RSLSVRMGKK

res gross greet greet with the first work work greet work greet was a second control of the cont

[Key to Table:]

Seq.IDNO	Name	Proteinsequenz
		VREYTMLRNI GLKTIDDIFK IAVHPLREAI DMRENMQNAI VSIKELCGLP PIASLKQCLL TLSSRLITSD
		NTPSVSLVMK DSFPYLEPLG AIPDVQKKML TAYDLMIQES RFLIEMADTV QEKIVQCQKA GMEFHEELHN
		LGAKEGLKGR KLNKATESFA WNITVLKGQG DLLKNAKNEA IENMKQIQLA CLSCGLSKAP SSSAEAKSKR
		SLEAIEEKES SEENGKL

THE STATE SHAME SH

[Key to Table:]